GenCore version 4.5

OM nucleic - nucleic search, using sw model

Run on: May 17, 2002, 23:38:58 ; Search time 139.1 Seconds
(without alignments)
1907.147 Million cell updates/sec

Title: US-09-719-748-1_COPY_62_1141
Perfect score: 1080
Sequence: 1 atggagccattcaagcagca.....ggaggaggagcagcacctcc 1080

Scoring table: IDENTITY_NUC
Gapop 10.0 , Gapext 1.0

Searched: 383533 seqs, 122816752 residues

Total number of hits satisfying chosen parameters: 767066

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : Issued_Patents_NA:*
1: /cgn2_6/ptodata/1/ina/5A_COMB.seq:*
2: /cgn2_6/ptodata/1/ina/5B_COMB.seq:*
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	% Query Match	Length	DB	ID	Description
1	514.6	47.6	2132	2	US-09-159-385-3	Sequence 3, Appli
2	514.6	47.6	2132	4	US-09-186-277-3	Sequence 3, Appli
3	512.8	47.5	1429	2	US-09-159-385-4	Sequence 4, Appli
4	512.8	47.5	1429	4	US-09-186-277-4	Sequence 4, Appli
5	448.6	41.5	4935	2	US-08-631-097-3	Sequence 3, Appli
6	448.6	41.5	5886	4	US-08-810-712-9	Sequence 9, Appli
7	445.4	41.2	480	3	US-09-221-235-12	Sequence 12, Appl
8	445.4	41.2	480	3	US-09-221-928-12	Sequence 12, Appl
9	445.4	41.2	480	3	US-09-221-527-12	Sequence 12, Appl
10	445.4	41.2	480	3	US-09-221-236-12	Sequence 12, Appl
11	445.4	41.2	480	3	US-09-221-416-12	Sequence 12, Appl
12	445.4	41.2	480	4	US-09-221-245-12	Sequence 12, Appl
13	445.4	41.2	480	4	US-09-163-115-12	Sequence 12, Appl
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15	445.4	41.2	480	4	US-09-593-553-12	Sequence 12, Appl
16	445.4	41.2	480	4	US-09-221-237-12	Sequence 12, Appl
17	445.4	41.2	1864	3	US-09-221-235-10	Sequence 10, Appl
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20	445.4	41.2	1864	3	US-09-221-236-10	Sequence 10, Appl
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28	146.8	13.6	1282	4	US-09-272-796-12	Sequence 12, Appl
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43	91.2	8.4	1349	3	US-08-459-595A-20	Sequence 20, Appl
44	91.2	8.4	1349	3	US-08-459-504B-20	Sequence 20, Appl
45	91.2	8.4	1349	3	US-08-459-444-20	Sequence 0, Appli

ALIGNMENTS

RESULT 1
US-09-159-385-3
; Sequence 3, Application US/09159385
; Patent No. 5958748
; GENERAL INFORMATION:
; APPLICANT: AKIRA, SHIZUO
; APPLICANT: KAWAI, TARO
; TITLE OF INVENTION: DNA CODING FOR SERINE/THREONINE KINASE
; FILE REFERENCE: PH-569
; CURRENT APPLICATION NUMBER: US/09/159,385
; CURRENT FILING DATE: 1998-09-23
; EARLIER APPLICATION NUMBER: JP97/261589
; EARLIER FILING DATE: 1997-09-26
; NUMBER OF SEQ ID NOS: 8
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 3
; LENGTH: 2132
; TYPE: DNA
; ORGANISM: Homo sapiens
; FEATURE:
; NAME/KEY: CDS
; LOCATION: (94)..(1455)
US-09-159-385-3

Query Match 47.6%; Score 514.6; DB 2; Length 2132;
Best Local Similarity 76.1%; Pred. No. 2.1e-125;
Matches 634; Conservative 0; Mismatches 199; Indels 0; Gaps 0;

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Qy     241 gacgtctatgagaaccgcaccgacgtggtgcacatccttgagctagtgtctggaggagag 300
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RESULT 2
US-09-186-277-3
; Sequence 3, Application US/09186277
; Patent No. 6171841
; GENERAL INFORMATION:
; APPLICANT: AKIRA, SHIZUO
; APPLICANT: KAWAI, TARO
; TITLE OF INVENTION: DNA CODING FOR SERINE/THREONINE KINASE
; FILE REFERENCE: 081356/0128
; CURRENT APPLICATION NUMBER: US/09/186,277
; CURRENT FILING DATE: 1998-11-05
; EARLIER APPLICATION NUMBER: JP97/261589
; EARLIER FILING DATE: 1997-09-26
; NUMBER OF SEQ ID NOS: 8
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 3
; LENGTH: 2132
; TYPE: DNA
; ORGANISM: Homo sapiens
; FEATURE:
; NAME/KEY: CDS
; LOCATION: (94)..(1455)
US-09-186-277-3

Query Match 47.6%; Score 514.6; DB 4; Length 2132;
Best Local Similarity 76.1%; Pred. No. 2.1e-125;
Matches 634; Conservative 0; Mismatches 199; Indels 0; Gaps 0;

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Db     94 atgtccacgttcaggcaggaggacgtggaggaccattatgagatgggggaggagctgggc 153

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Qy    121 gccaagttcatcaagaagcggcagagccgggcgagccggcgcggtgtgagccgggaggag 180
Db    214 gccaagttcatcaagaagcgccgcctgtcatccagccggcgtggggtgagccgggaggag 273

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Qy    301 ctcttcgatttcctggcccagaaggagtcactgagtgaggaggaggccaccagcttcatt 360
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Qy    361 aagcagatcctggatggggtgaactaccttcacacaaagaaaattgctcactttgatctc 420
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RESULT 3
US-09-159-385-4
; Sequence 4, Application US/09159385
; Patent No. 5958748
; GENERAL INFORMATION:
; APPLICANT: AKIRA, SHIZUO
; APPLICANT: KAWAI, TARO
; TITLE OF INVENTION: DNA CODING FOR SERINE/THREONINE KINASE
; FILE REFERENCE: PH-569
; CURRENT APPLICATION NUMBER: US/09/159,385
; CURRENT FILING DATE: 1998-09-23
; EARLIER APPLICATION NUMBER: JP97/261589
; EARLIER FILING DATE: 1997-09-26
; NUMBER OF SEQ ID NOS: 8
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 4
; LENGTH: 1429
; TYPE: DNA
; ORGANISM: Mus musculus
; FEATURE:
; NAME/KEY: CDS
; LOCATION: (10)..(1353)
US-09-159-385-4

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  Best Local Similarity   76.2%;  Pred. No. 5.3e-125;
  Matches  631;  Conservative    0;  Mismatches  197;  Indels    0;  Gaps    0;

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RESULT   4
US-09-186-277-4
; Sequence 4, Application US/09186277
; Patent No. 6171841
; GENERAL INFORMATION:
;  APPLICANT: AKIRA, SHIZUO
;  APPLICANT: KAWAI, TARO
;  TITLE OF INVENTION: DNA CODING FOR SERINE/THREONINE KINASE
;  FILE REFERENCE: 081356/0128
;  CURRENT APPLICATION NUMBER: US/09/186,277
;  CURRENT FILING DATE: 1998-11-05
;  EARLIER APPLICATION NUMBER: JP97/261589
;  EARLIER FILING DATE: 1997-09-26
;  NUMBER OF SEQ ID NOS: 8
;  SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 4
;   LENGTH: 1429
;   TYPE: DNA
;   ORGANISM: Mus musculus
;   FEATURE:
;   NAME/KEY: CDS
;   LOCATION: (10)..(1353)
US-09-186-277-4


  Query Match             47.5%;  Score 512.8;  DB 4;  Length 1429;
  Best Local Similarity   76.2%;  Pred. No. 5.3e-125;
  Matches  631;  Conservative    0;  Mismatches  197;  Indels    0;  Gaps    0;

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Qy     61 agtggccagtttgccatcgtgaagaagtgccgggagaagagcacggggcttgagtatgca 120
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Qy    121 gccaagttcatcaagaagcggcagagccgggcgagccggcgcggtgtgagccgggaggag 180
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Qy    661 acgaagcaggaaacactggcaaatatcacatcagtgagttacgactttgatgaggaattc 720
          || |||||||| || ||| | || ||| || ||||||  || ||||||||||||||| |
Db    670 accaagcaggagacgctgacgaacatctcagcagtgaactatgactttgatgaggaatac 729

Qy    721 ttcagccatacgagcgagctggccaaggactttattcggaagcttctggttaaagagacc 780
          ||||||   || |||||||||||||||||||| || || | ||| ||||| |||||  ||
Db    730 ttcagcagcaccagcgagctggccaaggacttcatccgcaggctgctggtcaaagacccc 789

Qy    781 cggaaacggctcacaatccaagaggctctcagacacccctggatcacg 828
            ||   || | || |||  | ||   ||    ||  |||||||| |
Db    790 aagaggaggatgaccatcgcacagagcctggagcattcctggatcaag 837
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RESULT   5
US-08-631-097-3
; Sequence 3, Application US/08631097
; Patent No. 5968816
;  GENERAL INFORMATION:
;    APPLICANT:  Kimchi, Adi
;    TITLE OF INVENTION:  Tumor Suppressor Genes,
;    TITLE OF INVENTION:  Proten Encloded Thereby, and Use of Said Genes and Protein
;    NUMBER OF SEQUENCES:  7
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  Wigman, Cohen, Leitner, & Myers, P.C.
;      STREET:  900 17th Street, N.W., Suite 1000
;      CITY:  Washington
;      STATE:  D.C.
;      COUNTRY:  USA
;      ZIP:  20006
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  Diskette, 3.50 inch, 1.44 Mb storage
;      COMPUTER:  IBM Compatible
;      OPERATING SYSTEM:  DOS
;      SOFTWARE:  ASCII
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/08/631,097
;      FILING DATE:  12-Apr-96
;      CLASSIFICATION:  514
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  PCT/US94/11598
;      FILING DATE:  12-Oct-94
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  Cohen, Herbert
;      REGISTRATION NUMBER:  25,109
;      REFERENCE/DOCKET NUMBER:  0744.057
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  (202)463-7700
;      TELEFAX:  (202)473-6915
;  INFORMATION FOR SEQ ID NO:  3:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  4935 base pairs
;      TYPE:  nucleic acid
;      STRANDEDNESS:  double
;      TOPOLOGY:  linear
;    MOLECULE TYPE:  Genomic DNA
;    HYPOTHETICAL:  No
;    ANTI-SENSE:  No
;    FRAGMENT TYPE:  No. 5968816 applicable
;    ORIGINAL SOURCE:
;      ORGANISM:  homo sapiens
;      STRAIN:  not applicable
;      INDIVIDUAL ISOLATE:  not applicable
;      DEVELOPMENTAL STAGE:  not applicable
;      HAPLOTYPE:  not applicable
;      TISSUE TYPE:  blood
;      CELL TYPE:  Leucocyte
;      CELL LINE:  HeLa
;      ORGANELLE:  not applicable
;    IMMEDIATE SOURCE:
;      LIBRARY:  not applicable
;      CLONE:  not applicable
;    POSITION IN GENOME:
;      CHROMOSOME/SEGMENT:  not applicable
;      MAP POSITION:  not applicable
;      UNITS:  not applicable
;    FEATURE:
;      NAME/KEY:  Seq. ID. NO.: 3 is
;      NAME/KEY:  the sequence in claim 1(iii) as Figure 8 of the specification
;      LOCATION:  not available
;      IDENTIFICATION METHOD:  experiment-
;      IDENTIFICATION METHOD:  in specification
;      OTHER INFORMATION:  prevention of IFN-2
;      OTHER INFORMATION:  promoted cell death
;    PUBLICATION INFORMATION:  not available
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US-08-631-097-3

Query Match 41.5%; Score 448.6; DB 2; Length 4935;
Best Local Similarity 67.5%; Pred. No. 5.5e-108;
Matches 659; Conservative 0; Mismatches 294; Indels 24; Gaps 1;

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Qy      10 ttcaagcagcagaaggtggaggacttttatgacatcggagaggagctggggagtggccag 69
           |||| |||| | || ||||| || |  || |||| ||| ||||| || || ||||| |||
Db     346 TTCAGGCAGGAAAACGTGGATGATTACTACGACACCGGCGAGGAACTTGGCAGTGGACAG 405

Qy      70 tttgccatcgtgaagaagtgccgggagaagagcacggggcttgagtatgcagccaagttc 129
           |||||  | |||||||| ||||| ||||| || || || ||  ||||| | ||||| |||
Db     406 TTTGCGGTTGTGAAGAAATGCCGTGAGAAAAGTACCGGCCTCCAGTATCCCGCCAAATTC 465

Qy     130 atcaagaagcggcagagccgggcgagccggcgcggtgtgagccgggaggagatcgagcgg 189
           |||||||| ||  ||    | | |||||||| |||||||||||| ||||| |||||||||
Db     466 ATCAAGAAAAGGAGGACTAAGTCCAGCCGGCGGGGTGTGAGCCGCGAGGACATCGAGCGG 525

Qy     190 gaggtgagcatcctgcggcaggtgctgcaccacaatgtcatcacgctgcacgacgtctat 249
           |||||  |||||||||  | || | | ||||| |||||||||||| |||||||| ||||||
Db     526 GAGGTCAGCATCCTGAAGGAGATCCAGCACCCCAATGTCATCACCCTGCACGAGGTCTAT 585

Qy     250 gagaaccgcaccgacgtggtgcacatccttgagctagtgtctggaggagagctcttcgat 309
           ||||||   || |||||  | |  ||| | || || ||  | || || ||||| || ||
Db     586 GAGAACAAGACGGACGTCATCCTGATCTTGGAACTCGTTGCAGGTGGCGAGCTGTTTGAC 645

Qy     310 ttcctggcccagaaggagtcactgagtgaggaggaggccaccagcttcattaagcagatc 369
           ||| | ||  | ||||| ||  | | ||| ||||| || ||    ||  | || || ||
Db     646 TTCTTAGCTGAAAAGGAATCTTTAACTGAAGAGGAAGCAACTGAATTTCTCAAACAAATT 705

Qy     370 ctggatggggtgaactaccttcacacaaagaaaattgctcactttgatctcaagccagaa 429
           ||  |||| ||  ||||||| ||| |     |||| || |||||||||| ||||| ||
Db     706 CTTAATGGTGTTTACTACCTGCACTCCCTTCAAATCGCCCACTTTGATCTTAAGCCTGAG 765

Qy     430 aacattatgttgttagacaagaatattcccattccacacatcaagctgattgactttggt 489
           ||||| ||| | || || |  ||| | ||||  || |  |||||| | ||||||||
Db     766 AACATAATGCTTTTGGATAGAAATGTCCCCAAACCTCGGATCAAGATCATTGACTT---- 821

Qy     490 ctggctcacgaaatagaagatggagttgaatttaagaatatttttgggacgccggaattt 549
                                |||| |||||||||| || || |||||||| || || |||
Db     822 --------------------TGGAAATGAATTTAAAAACATATTTGGGACTCCAGAGTTT 861

Qy     550 gttgctccagaaattgtgaactacgagcccctgggtctggaggctgacatgtggagcata 609
           || ||||| || || || ||||| || || || ||||| ||||| || |||||||| ||
Db     862 GTCGCTCCTGAGATAGTCAACTATGAACCTCTTGGTCTTGAGGCAGATATGTGGAGTATC 921

Qy     610 ggcgtcatcacctacatcctcttaagtggagcatccccttttcctgggagacacgaagcag 669
           || || || ||||| |||||| ||||||| || ||||| || || |||||||| |||||
Db     922 GGGGTAATAACCTATATCCTCCTAAGTGGGGCCTCCCCATTTCTTGGAGACACTAAGCAA 981

Qy     670 gaaacactggcaaatatcacatcagtgagttacgactttgatgaggaattcttcagccat 729
           |||||  | |||||| |  |  | || |  ||||| ||||| || |||| ||||||  ||
Db     982 GAAACGTTAGCAAATGTATCCGCTGTCAACTACGAATTTGAGGATGAATACTTCAGTAAT 1041

Qy     730 acgagcgagctggccaaggactttattcggaagcttctggttaaagagacccggaaacgg 789
           || || |  || ||||| || || ||  | |  |||||||| || ||  |   |||  |
Db    1042 ACCAGTGCCCTAGCCAAAGATTTCATAAGAAGACTTCTGGTCAAGGATCCAAAGAAGAGA 1101

Qy     790 ctcacaatccaagaggctctcagacacccctggatcacgccggtggacaaccagcaagcc 849
            | ||||| |||||   | |    || ||||||||||| |||    || |  || || ||
Db    1102 ATGACAATTCAAGATAGTTTGCAGCATCCCTGGATCAAGCCTAAAGATACACAACAGGCA 1161

Qy     850 atggtgcgacgggagtctgtggtcaatctggagaacttcaggaagcagtatgtccgcagg 909
           |     ||   | || |  || ||  ||||||| |||| ||||       | ||| |
Db    1162 CTTAGTAGAAAAGCATCAGCAGTAAACATGGAGAAATTCAAGAAGTTTGCAGCCCGGAAA 1221

Qy     910 cggtggaagctttccttcagcatcgtgtccctgtgcaaccacctcacccgctcgctgatg 969
           |||||  |  |||  |  || |  | || ||||||  |     |  || |  ||  |  ||
Db    1222 AAATGGAAACAATCCGTTCGCTTGATATCACTGTGCCAAAGATTATCCAGGTCATTCCTG 1281

Qy     970 aagaaggtgcacctgag 986
              |      || ||||
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Db   1282 TCCAGAAGTAACATGAG 1298


RESULT   6
US-08-810-712-9
; Sequence 9, Application US/08810712G
; Patent No. 6160106
; GENERAL INFORMATION:
;  APPLICANT: Yeda Research and Development  Co. LTD
;  TITLE OF INVENTION: Tumor Suppressor Genes, Proteins Encoded Thereby and
;  TITLE OF INVENTION: Use of said Genes and Proteins
;  FILE REFERENCE: sequencelist
;  CURRENT APPLICATION NUMBER: US/08/810,712G
;  CURRENT FILING DATE: 1997-03-03
;  EARLIER APPLICATION NUMBER: PCT/US94/11598
;  EARLIER FILING DATE: 1994-10-12
;  NUMBER OF SEQ ID NOS: 31
;  SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 9
;   LENGTH: 5886
;   TYPE: DNA
;   ORGANISM: Homo sapiens
;   FEATURE:
;   NAME/KEY: CDS
;   LOCATION: (337)..(4605)
US-08-810-712-9


  Query Match             41.5%;  Score 448.6;  DB 4;  Length 5886;
  Best Local Similarity   67.5%;  Pred. No. 5.9e-108;
  Matches  659;  Conservative    0;  Mismatches  294;  Indels   24;  Gaps    1;

Qy      10 ttcaagcagcagaaggtggaggacttttatgacatcggagaggagctggggagtggccag 69
           |||| |||| | || ||||| || |  || |||| ||| ||||| || || ||||| |||
Db     346 ttcaggcaggaaaacgtggatgattactacgacaccggcgaggaacttggcagtggacag 405

Qy      70 tttgccatcgtgaagaagtgccgggagaagagcacggggcttgagtatgcagccaagttc 129
           |||||  | |||||||| ||||| ||||| || || || ||  ||||| | ||||| |||
Db     406 tttgcggttgtgaagaaatgccgtgagaaaagtaccggcctccagtatcccgccaaattc 465

Qy     130 atcaagaagcggcagagccgggcgagccggcgcggtgtgagccgggaggagatcgagcgg 189
           ||||||||  ||  ||    | | |||||||| |||||||||| ||||| |||||||||
Db     466 atcaagaaaaggaggactaagtccagccggcggggtgtgagccgcgaggacatcgagcgg 525

Qy     190 gaggtgagcatcctgcggcaggtgctgcaccacaatgtcatcacgctgcacgacgtctat 249
           ||||| |||||||||  | || | | ||||| |||||||||||| |||||||| ||||||
Db     526 gaggtcagcatcctgaaggagatccagcaccccaatgtcatcaccctgcacgaggtctat 585

Qy     250 gagaaccgcaccgacgtggtgcacatccttgagctagtgtctggaggagagctcttcgat 309
           ||||||   || |||||  | |  ||| | || || ||  | || || ||||| || ||
Db     586 gagaacaagacggacgtcatcctgatcttggaactcgttgcaggtggcgagctgtttgac 645

Qy     310 ttcctggcccagaaggagtcactgagtgaggaggaggccaccagcttcattaagcagatc 369
           ||| | ||  | ||||| ||  | | ||| ||||| || ||    ||  | || || ||
Db     646 ttcttagctgaaaaggaatctttaactgaagaggaagcaactgaatttctcaaacaaatt 705

Qy     370 ctggatggggtgaactaccttcacacaaagaaaattgctcactttgatctcaagccagaa 429
           ||  |||| ||  ||||||| ||| |     |||| || |||||||||| ||||| ||
Db     706 cttaatggtgtttactacctgcactcccttcaaatcgcccactttgatcttaagcctgag 765

Qy     430 aacattatgttgttagacaagaatattcccattccacacatcaagctgattgactttggt 489
           ||||| ||| | || || |  ||| | |||  || |  |||||| | ||||||||
Db     766 aacataatgcttttggatagaaatgtccccaaacctcggatcaagatcattgactt---- 821

Qy     490 ctggctcacgaaatagaagatggagttgaatttaagaatatttttgggacgccggaattt 549
                           ||||  |||||||||| || || |||||||| || || |||
Db     822 --------------------tggaaatgaatttaaaaacatatttgggactccagagttt 861

Qy     550 gttgctccagaaattgtgaactacgagcccctgggtctggaggctgacatgtggagcata 609
           || ||||| || || || ||||| || || || ||||| ||||| || |||||||| ||
Db     862 gtcgctcctgagatagtcaactatgaacctcttggtcttgaggcagatatgtggagtatc 921

Qy     610 ggcgtcatcacctacatcctcttaagtggagcatccccttttcctgggagacacgaagcag 669
           || || || ||||| |||||| ||||||| || ||||| || || |||||||| |||||
Db     922 ggggtaataacctatatcctcctaagtggggcctccccatttcttggagacactaagcaa 981

Qy     670 gaaacactggcaaatatcacatcagtgagttacgactttgatgaggaattcttcagccat 729
           |||||  | |||||| |  |  | || |  ||||| ||||| || |||| ||||||  ||
Db     982 gaaacgttagcaaatgtatccgctgtcaactacgaatttgaggatgaatacttcagtaat 1041

Qy     730 acgagcgagctggccaaggactttattcggaagcttctggttaaagagacccggaaacgg 789
           || || |  || ||||| || || ||  | |  |||||||| || ||  |   |||  |
Db    1042 accagtgccctagccaaagatttcataagaagacttctggtcaaggatccaaagaagaga 1101

Qy     790 ctcacaatccaagaggctctcagacacccctggatcacgccggtggacaaccagcaagcc 849
           | ||||| |||||   | |    || ||||||||||| |||    || |  || || ||
Db    1102 atgacaattcaagatagtttgcagcatccctggatcaagcctaaagatacacaacaggca 1161

Qy     850 atggtgcgacgggagtctgtggtcaatctggagaacttcaggaagcagtatgtccgcagg 909
           |     ||   |  || |  || ||  ||||||| |||| ||||      | ||| |
Db    1162 cttagtagaaaagcatcagcagtaaacatggagaaattcaagaagtttgcagcccggaaa 1221

Qy     910 cggtggaagctttccttcagcatcgtgtccctgtgcaaccacctcacccgctcgctgatg 969
           |||||  |  ||| |  || |  | || |||||| |     |  || | ||  |  ||
Db    1222 aaatggaaacaatccgttcgcttgatatcactgtgccaaagattatccaggtcattcctg 1281

Qy     970 aagaaggtgcacctgag 986
           |      || ||||
Db    1282 tccagaagtaacatgag 1298


RESULT   7
US-09-221-235-12
; Sequence 12, Application US/09221235
; Patent No. 6043040
; GENERAL INFORMATION:
;  APPLICANT: Acton, Susan
;  TITLE OF INVENTION: NOVEL CSAPK-1 NUCLEIC ACID MOLECULES AND USES THEREFOR
;  FILE REFERENCE: MNI-050
;  CURRENT APPLICATION NUMBER: US/09/221,235
;  CURRENT FILING DATE: 1998-12-28
;  EARLIER APPLICATION NUMBER: 09/163,115
;  EARLIER FILING DATE:
;  NUMBER OF SEQ ID NOS: 15
;  SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 12
;   LENGTH: 480
;   TYPE: DNA
;   ORGANISM: Homo sapiens
;   FEATURE:
;   NAME/KEY: CDS
;   LOCATION: (1)..(480)
US-09-221-235-12


  Query Match             41.2%;  Score 445.4;  DB 3;  Length 480;
  Best Local Similarity   98.7%;  Pred. No. 1.5e-107;
  Matches  449;  Conservative    0;  Mismatches    6;  Indels    0;  Gaps    0;

Qy     626 tcctcttaagtggagcatccccttttcctgggagacacgaagcaggaaacactggcaaata 685
           ||  ||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      26 tcagcttaagtggagcatccccttttcctgggagacacgaagcaggaaacactggcaaata 85

Qy     686 tcacatcagtgagttacgactttgatgaggaattcttcagccatacgagcgagctggcca 745
           ||||| |||||||||||||||||||||||||||||||||||| ||||||||||||||||
Db      86 tcacagcagtgagttacgactttgatgaggaattcttcagccagacgagcgagctggcca 145

Qy     746 aggactttattcggaagcttctggttaaagagacccggaaacggctcacaatccaagagg 805
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     146 aggactttattcggaagcttctggttaaagagacccggaaacggctcacaatccaagagg 205

Qy     806 ctctcagacacccctggatcacgccggtggacaaccagcaagccatggtgcgacgggagt 865
           |||||||||||||||||||||||||||||||||||||||||||||||||||  ||||||
Db     206 ctctcagacacccctggatcacgccggtggacaaccagcaagccatggtgcgcagggagt 265

Qy     866 ctgtggtcaatctggagaacttcaggaagcagtatgtccgcaggcggtggaagctttcct 925
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Db    266 ctgtggtcaatctggagaacttcaggaagcagtatgtccgcaggcggtggaagctttcct 325

Qy    926 tcagcatcgtgtccctgtgcaaccacctcacccgctcgctgatgaagaaggtgcacctga 985
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    326 tcagcatcgtgtccctgtgcaaccacctcacccgctcgctgatgaagaaggtgcacctga 385

Qy    986 ggccggatgaggacctgaggaactgtgagagtgacactgaggaggacatcgccaggagga 1045
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    386 ggccggatgaggacctgaggaactgtgagagtgacactgaggaggacatcgccaggagga 445

Qy   1046 aagccctccacccacggaggaggagcagcacctcc 1080
          |||||||||||||||||||||||||||||||||||
Db    446 aagccctccacccacggaggaggagcagcacctcc 480
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RESULT 8
US-09-221-928-12
; Sequence 12, Application US/09221928
; Patent No. 6121030
; GENERAL INFORMATION:
; APPLICANT: Acton, Susan
; TITLE OF INVENTION: NOVEL CSAPK-1 NUCLEIC ACID MOLECULES AND USES THEREFOR
; FILE REFERENCE: MNI-050
; CURRENT APPLICATION NUMBER: US/09/221,928
; CURRENT FILING DATE: 1998-12-28
; EARLIER APPLICATION NUMBER: 09/163,115
; EARLIER FILING DATE:
; NUMBER OF SEQ ID NOS: 15
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 12
; LENGTH: 480
; TYPE: DNA
; ORGANISM: Homo sapiens
; FEATURE:
; NAME/KEY: CDS
; LOCATION: (1)..(480)
US-09-221-928-12

Query Match 41.2%; Score 445.4; DB 3; Length 480;
Best Local Similarity 98.7%; Pred. No. 1.5e-107;
Matches 449; Conservative 0; Mismatches 6; Indels 0; Gaps 0;

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Qy    626 tcctcttaagtggagcatccccctttcctgggagacacgaagcaggaaacactggcaaata 685
          ||  ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     26 tcagcttaagtggagcatccccctttcctgggagacacgaagcaggaaacactggcaaata 85

Qy    686 tcacatcagtgagttacgactttgatgaggaattcttcagccatacgagcgagctggcca 745
          |||||  ||||||||||||||||||||||||||||||||| ||||||||||||||||||
Db     86 tcacagcagtgagttacgactttgatgaggaattcttcagccagacgagcgagctggcca 145

Qy    746 aggactttattcggaagcttctggttaaagagacccggaaacggctcacaatccaagagg 805
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    146 aggactttattcggaagcttctggttaaagagacccggaaacggctcacaatccaagagg 205

Qy    806 ctctcagacacccctggatcacgccggtggacaaccagcaagccatggtgcgacgggagt 865
          ||||||||||||||||||||||||||||||||||||||||||||||||||  ||||||
Db    206 ctctcagacacccctggatcacgccggtggacaaccagcaagccatggtgcgcagggagt 265

Qy    866 ctgtggtcaatctggagaacttcaggaagcagtatgtccgcaggcggtggaagctttcct 925
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    266 ctgtggtcaatctggagaacttcaggaagcagtatgtccgcaggcggtggaagctttcct 325

Qy    926 tcagcatcgtgtccctgtgcaaccacctcacccgctcgctgatgaagaaggtgcacctga 985
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    326 tcagcatcgtgtccctgtgcaaccacctcacccgctcgctgatgaagaaggtgcacctga 385

Qy    986 ggccggatgaggacctgaggaactgtgagagtgacactgaggaggacatcgccaggagga 1045
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    386 ggccggatgaggacctgaggaactgtgagagtgacactgaggaggacatcgccaggagga 445

Qy   1046 aagccctccacccacggaggaggagcagcacctcc 1080
          |||||||||||||||||||||||||||||||||||
Db    446 aagccctccacccacggaggaggagcagcacctcc 480
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RESULT 9
US-09-221-527-12
; Sequence 12, Application US/09221527
; Patent No. 6146832
; GENERAL INFORMATION:
; APPLICANT: Acton, Susan
; TITLE OF INVENTION: NOVEL CSAPK-1 NUCLEIC ACID MOLECULES AND USES THEREFOR
; FILE REFERENCE: MNI-050
; CURRENT APPLICATION NUMBER: US/09/221,527
; CURRENT FILING DATE: 1998-12-28
; EARLIER APPLICATION NUMBER: 09/163,115
; EARLIER FILING DATE:
; NUMBER OF SEQ ID NOS: 15
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 12
; LENGTH: 480
; TYPE: DNA
; ORGANISM: Homo sapiens
; FEATURE:
; NAME/KEY: CDS
; LOCATION: (1)..(480)
US-09-221-527-12

Query Match 41.2%; Score 445.4; DB 3; Length 480;
Best Local Similarity 98.7%; Pred. No. 1.5e-107;
Matches 449; Conservative 0; Mismatches 6; Indels 0; Gaps 0;

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          ||  ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     26 tcagcttaagtggagcatccccctttcctgggagacacgaagcaggaaacactggcaaata 85

Qy    686 tcacatcagtgagttacgactttgatgaggaattcttcagccatacgagcgagctggcca 745
          |||||  ||||||||||||||||||||||||||||||||| ||||||||||||||||||
Db     86 tcacagcagtgagttacgactttgatgaggaattcttcagccagacgagcgagctggcca 145

Qy    746 aggactttattcggaagcttctggttaaagagacccggaaacggctcacaatccaagagg 805
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    146 aggactttattcggaagcttctggttaaagagacccggaaacggctcacaatccaagagg 205

Qy    806 ctctcagacacccctggatcacgccggtggacaaccagcaagccatggtgcgacgggagt 865
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Qy    866 ctgtggtcaatctggagaacttcaggaagcagtatgtccgcaggcggtggaagctttcct 925
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Qy    926 tcagcatcgtgtccctgtgcaaccacctcacccgctcgctgatgaagaaggtgcacctga 985
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Qy    986 ggccggatgaggacctgaggaactgtgagagtgacactgaggaggacatcgccaggagga 1045
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RESULT 10
US-09-221-236-12
; Sequence 12, Application US/09221236
; Patent No. 6146841
; GENERAL INFORMATION:
; APPLICANT: Acton, Susan
; TITLE OF INVENTION: NOVEL CSAPK-1 NUCLEIC ACID MOLECULES AND USES THEREFOR
; FILE REFERENCE: MNI-050

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;  CURRENT APPLICATION NUMBER: US/09/221,236
;  CURRENT FILING DATE: 1998-12-28
;  EARLIER APPLICATION NUMBER: 09/163,115
;  EARLIER FILING DATE: 1998-09-29
;  NUMBER OF SEQ ID NOS: 15
;  SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 12
;   LENGTH: 480
;   TYPE: DNA
;   ORGANISM: Homo sapiens
;   FEATURE:
;   NAME/KEY: CDS
;   LOCATION: (1)..(480)
US-09-221-236-12


  Query Match             41.2%;  Score 445.4;  DB 3;  Length 480;
  Best Local Similarity   98.7%;  Pred. No. 1.5e-107;
  Matches  449;  Conservative    0;  Mismatches    6;  Indels    0;  Gaps    0;

Qy    626 tcctcttaagtggagcatccccttcctgggagacacgaagcaggaaacactggcaaata 685
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Qy    686 tcacatcagtgagttacgactttgatgaggaattcttcagccatacgagcgagctggcca 745
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Db     86 tcacagcagtgagttacgactttgatgaggaattcttcagccagacgagcgagctggcca 145

Qy    746 aggactttattcggaagcttctggttaaagagacccggaaacggctcacaatccaagagg 805
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Db    146 aggactttattcggaagcttctggttaaagagacccggaaacggctcacaatccaagagg 205

Qy    806 ctctcagacacccctggatcacgccggtggacaaccagcaagccatggtgcgacgggagt 865
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Qy    866 ctgtggtcaatctggagaacttcaggaagcagtatgtccgcaggcggtggaagctttcct 925
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Qy    926 tcagcatcgtgtccctgtgcaaccacctcacccgctcgctgatgaagaaggtgcacctga 985
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Qy    986 ggccggatgaggacctgaggaactgtgagagtgacactgaggaggacatcgccaggagga 1045
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Qy   1046 aagccctccacccacggaggaggagcagcacctcc 1080
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Db    446 aagccctccacccacggaggaggagcagcacctcc 480


RESULT  11
US-09-221-416-12
; Sequence 12, Application US/09221416
; Patent No. 6153417
; GENERAL INFORMATION:
;  APPLICANT: Acton, Susan
;  TITLE OF INVENTION: NOVEL CSAPK-1 NUCLEIC ACID MOLECULES AND USES THEREFOR
;  FILE REFERENCE: MNI-050
;  CURRENT APPLICATION NUMBER: US/09/221,416
;  CURRENT FILING DATE: 1998-12-28
;  EARLIER APPLICATION NUMBER: 09/163,115
;  EARLIER FILING DATE: 1998-09-29
;  NUMBER OF SEQ ID NOS: 15
;  SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 12
;   LENGTH: 480
;   TYPE: DNA
;   ORGANISM: Homo sapiens
;   FEATURE:
;   NAME/KEY: CDS
;   LOCATION: (1)..(480)
US-09-221-416-12


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Qy    686 tcacatcagtgagttacgactttgatgaggaattcttcagccatacgagcgagctggcca 745
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Qy    746 aggactttattcggaagcttctggttaaagagacccggaaacggctcacaatccaagagg 805
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Qy    866 ctgtggtcaatctggagaacttcaggaagcagtatgtccgcaggcggtggaagctttcct 925
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Qy    926 tcagcatcgtgtccctgtgcaaccacctcacccgctcgctgatgaagaaggtgcacctga 985
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Qy    986 ggccggatgaggacctgaggaactgtgagagtgacactgaggaggacatcgccaggagga 1045
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RESULT  12
US-09-221-245-12
; Sequence 12, Application US/09221245
; Patent No. 6180358
; GENERAL INFORMATION:
;  APPLICANT: Acton, Susan
;  TITLE OF INVENTION: NOVEL CSAPK-1 NUCLEIC ACID MOLECULES AND USES THEREFOR
;  FILE REFERENCE: MNI-050
;  CURRENT APPLICATION NUMBER: US/09/221,245
;  CURRENT FILING DATE: 1998-12-28
;  EARLIER APPLICATION NUMBER: US 09/163,115
;  EARLIER FILING DATE: 1998-09-29
;  NUMBER OF SEQ ID NOS: 15
;  SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 12
;   LENGTH: 480
;   TYPE: DNA
;   ORGANISM: Homo sapiens
;   FEATURE:
;   NAME/KEY: CDS
;   LOCATION: (1)..(480)
US-09-221-245-12


  Query Match             41.2%;  Score 445.4;  DB 4;  Length 480;
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Qy    686 tcacatcagtgagttacgactttgatgaggaattcttcagccatacgagcgagctggcca 745
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Db     86 tcacagcagtgagttacgactttgatgaggaattcttcagccagacgagcgagctggcca 145

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Db    146 aggactttattcggaagcttctggttaaagagacccggaaacggctcacaatccaagagg 205

Qy    806 ctctcagacacccctggatcacgccggtggacaaccagcaagccatggtgcgacgggagt 865
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Qy    866 ctgtggtcaatctggagaacttcaggaagcagtatgtccgcaggcggtggaagctttcct 925
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Qy    926 tcagcatcgtgtccctgtgcaaccacctcacccgctcgctgatgaagaaggtgcacctga 985
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Qy    986 ggccggatgaggacctgaggaactgtgagagtgacactgaggaggacatcgccaggagga 1045
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RESULT  13
US-09-163-115-12
; Sequence 12, Application US/09163115A
; Patent No. 6183962
; GENERAL INFORMATION:
;  APPLICANT: Acton, Susan
;  TITLE OF INVENTION: NOVEL CSAPK-1 NUCLEIC ACID MOLECULES AND USES THEREFOR
;  FILE REFERENCE: MNI-050
;  CURRENT APPLICATION NUMBER: US/09/163,115A
;  CURRENT FILING DATE: 1998-09-29
;  NUMBER OF SEQ ID NOS: 15
;  SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 12
;   LENGTH: 480
;   TYPE: DNA
;   ORGANISM: Homo sapiens
;   FEATURE:
;   NAME/KEY: CDS
;   LOCATION: (1)..(480)
US-09-163-115-12


  Query Match             41.2%;  Score 445.4;  DB 4;  Length 480;
  Best Local Similarity   98.7%;  Pred. No. 1.5e-107;
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Qy    686 tcacatcagtgagttacgactttgatgaggaattcttcagccatacgagcgagctggcca 745
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Db     86 tcacagcagtgagttacgactttgatgaggaattcttcagccagacgagcgagctggcca 145

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Qy    926 tcagcatcgtgtccctgtgcaaccacctcacccgctcgctgatgaagaaggtgcacctga 985
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Qy    986 ggccggatgaggacctgaggaactgtgagagtgacactgaggaggacatcgccaggagga 1045
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RESULT  14
US-09-221-528-12
; Sequence 12, Application US/09221528
; Patent No. 6190874
; GENERAL INFORMATION:
;  APPLICANT: Acton, Susan
;  TITLE OF INVENTION: NOVEL CSAPK-1 NUCLEIC ACID MOLECULES AND USES THEREFOR
;  FILE REFERENCE: MNI-050
;  CURRENT APPLICATION NUMBER: US/09/221,528
;  CURRENT FILING DATE: 1998-12-28
;  EARLIER APPLICATION NUMBER: 09/163,115
;  EARLIER FILING DATE: 1998-09-29
;  NUMBER OF SEQ ID NOS: 15
;  SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 12
;   LENGTH: 480
;   TYPE: DNA
;   ORGANISM: Homo sapiens
;   FEATURE:
;   NAME/KEY: CDS
;   LOCATION: (1)..(480)
US-09-221-528-12


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Qy    686 tcacatcagtgagttacgactttgatgaggaattcttcagccatacgagcgagctggcca 745
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Qy    926 tcagcatcgtgtccctgtgcaaccacctcacccgctcgctgatgaagaaggtgcacctga 985
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RESULT  15
US-09-593-553-12
; Sequence 12, Application US/09593553
; Patent No. 6200770
; GENERAL INFORMATION:
;  APPLICANT:  Acton, Susan
;  TITLE OF INVENTION:  NOVEL CSAPK-1 NUCLEIC ACID MOLECULES AND USES THEREFOR
;  FILE REFERENCE:  MNI-050
;  CURRENT APPLICATION NUMBER: US/09/593,553
;  CURRENT FILING DATE:  2000-06-14
;  PRIOR APPLICATION NUMBER:  09/163,115
;  PRIOR FILING DATE:  1998-09-28
;  NUMBER OF SEQ ID NOS:  15
;  SOFTWARE:  PatentIn Ver. 2.0
; SEQ ID NO 12
;   LENGTH: 480
;   TYPE: DNA
;   ORGANISM: Homo sapiens
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;   NAME/KEY: CDS
;   LOCATION: (1)..(480)
US-09-593-553-12


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Qy    686 tcacatcagtgagttacgactttgatgaggaattcttcagccatacgagcgagctggcca 745
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Qy    866 ctgtggtcaatctggagaacttcaggaagcagtatgtccgcaggcggtggaagctttcct 925
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Qy    926 tcagcatcgtgtccctgtgcaaccacctcacccgctcgctgatgaagaaggtgcacctga 985
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Qy    986 ggccggatgaggacctgaggaactgtgagagtgacactgaggaggacatcgccaggagga 1045
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Db    386 ggccggatgaggacctgaggaactgtgagagtgacactgaggaggacatcgccaggagga 445

Qy   1046 aagccctccacccacggaggaggagcagcacctcc 1080
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